

1 Phytoplankton as a principal diet for callianassid shrimp larvae in coastal
2 waters, estimated from laboratory rearing and stable isotope analysis

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16 ABSTRACT: The field diet of meroplanktonic decapod crustacean larvae is poorly known,
17 despite standard use of microzooplankton as food in laboratory culture. Using callianassid
18 shrimp *Nihonotrypaea harmandi* larvae collected from a 65 m deep inner-shelf location off
19 mid-western Kyushu, Japan, between June and August 2012 and 2013 and mass-reared in the
20 laboratory, phytoplankton-based diet through larval development (zoeae I–VI to decapodid)
21 was demonstrated. When the pure-cultured diatom *Chaetoceros gracilis* was fed to zoeae,
22 survival rate to decapodids was 3.4 to 3.9% in 26 to 40 d at 22°C, which was comparable to
23 previous rearing results for zoeae fed microzooplankton. Trophic enrichment factors (TEFs)

24 from stable isotope (SI) analysis of zoeal whole-body tissue in the laboratory were 2.0‰ for
25 $\delta^{13}\text{C}$ and 1.9‰ for $\delta^{15}\text{N}$. In the field water column, diatoms dominated the nano- to micro-sized
26 plankton, accounting for 38 to 81% of the biovolume, followed by heterotrophic protists. The
27 trophic position (TP) estimated from amino acid-specific $\delta^{15}\text{N}$ values for the field-collected
28 zoeae VI was 2.1 ($\text{TP}_{\text{Glu/Phe}}$) or 2.7 ($\text{TP}_{\text{Ala/Phe}}$), suggesting that those zoeae fed on mixtures of
29 phytoplankton and heterotrophs including protists. Bulk SI analyses were performed for
30 particulate organic matter (POM; proxy for phytoplankton), microzooplankton (mainly
31 calanoid copepods), and shrimp zoeae to elucidate the diet of larvae in the water column. A
32 shift in SI from fresh to degraded POM was determined through the incubation of
33 field-collected POM. Based on this shift during degradation and larval TEFs, phytoplankton
34 and their sinking detritus with heterotrophic protists were estimated to be the principal diet for
35 those larvae residing mostly below the chlorophyll maximum layer.

36 KEY WORDS: *Nihonotrypaea harmandi* · Zoea · Rearing · Diatom · Isotopic trophic
37 enrichment factor · Amino-acid- $\delta^{15}\text{N}$ -based trophic position · Phytodetritus · Protists

38 INTRODUCTION

39 Meroplanktonic larvae of macrobenthos are a seasonally significant component of
40 zooplankton assemblages in tropical, temperate, and boreal estuarine and coastal waters. The
41 successful acquisition of food by planktotrophic larvae is one critical process in their normal
42 development and survival (Thorson 1950, Day & McEdward 1984, Olson & Olson 1989,
43 Metaxas & Saunders 2009). Decapod crustaceans are a common component of both benthic
44 macro-invertebrate communities and meroplanktonic assemblages in these nearshore waters.
45 The potential food sources for planktotrophic decapod larvae comprise bacteria, phytoplankton
46 ranging from pico- to micro-sized groups, zooplankton ranging from nano- to meso-sized
47 groups, and detritus and fecal pellets (Lebour 1922, Stickney & Perkins 1981, Factor & Dexter
48 1993, Mascetti & Wehrtmann 1996, Lehto et al. 1998, Anger 2001, Schwamborn et al. 2006,

49 Fileman et al. 2014). In laboratory rearing of decapod larvae, brine shrimp *Artemia* spp. nauplii
50 and rotifers (e.g. *Brachionus* spp.), which are microzooplankton never encountered by those
51 larvae in their natural environment, have most frequently been used for complete larval
52 development from newly hatched to post-larval stages (McConaugha 1985, Harvey & Epifanio
53 1997, Anger 2001, Calado et al. 2010). Such success in larval culture, often with excess supply
54 of those food items, does not necessarily mean that microzooplankton are a principal food
55 source for natural larval populations, as individual density, microscale patchiness, filterability
56 (e.g. non-motile phytoplankton by filter-feeding larvae) or catchability (e.g. motile
57 phytoplankton or zooplankton by raptorial larvae), digestibility, and nutritional value are likely
58 to play their roles in natural diets. Principal diets of decapod crustacean larvae under natural
59 conditions thus remain to be determined.

60 Successful rearing of planktotrophic larvae of species in 1 of the 2 decapod suborders,
61 Pleocyemata, to the decapodid (= post-larval) or juvenile stage solely using phytoplankton has
62 been achieved in only a few studies on brachyuran crabs (Atkins 1955, Bousquette 1980,
63 Harms & Seeger 1989). In the latter study, larvae of a majid fed a diatom species showed a
64 longer developmental duration and lower survival rate than those fed *Artemia* sp. nauplii. In
65 other studies, phytoplankton are regarded as partially valid foods, typically compared with 2
66 control rearing treatments, either fed *Artemia* spp. nauplii (complete) or starved (invalid)
67 (Table 1). For zoeal stages in penaeid shrimp (suborder Dendrobranchiata), phytoplankton
68 (mainly diatoms) are considered a fully effective food source (Preston et al. 1992, Yano 2005).

69 Even if phytoplankton are a principal diet of meroplanktonic decapod crustacean
70 larvae to complete their development in the field, examination alone of ingested food species
71 contained in larval digestive tracts or fecal pellets, including algal cell walls, photosynthetic
72 pigments, and DNA-identified amorphous material, will not suffice to reach a convincing
73 conclusion. In addition to live forms, phytodetritus can be a potential food source for decapod

74 larvae (Meyer-Harms & Harms 1993, Kiørboe 2011, Turner 2015). Although carbon and
75 nitrogen stable isotope ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$) analysis is one promising way to estimate diets of those
76 larvae, the number of actual isotope studies is scarce. In a study area ranging from a mangrove
77 shore to an offshore shelf in northeastern Brazil, the contribution of mangrove carbon to
78 decapod larval nutrition was regarded as negligible, and pelagic primary producers were
79 suggested as principal diet sources (Schwamborn et al. 1999, 2002). However, the estimation
80 of the trophic position (TP) and food sources for target species based on such ‘bulk’ isotope
81 analysis has uncertainties. First, it is impossible to cover all potential food sources in the field,
82 where a variety of primary producers and consumers exist. Second, isotope values of primary
83 producers are variable in space and time due to the variability of their species composition, and
84 dissolved inorganic carbon and nitrogen sources. Finally, as enrichment of ^{13}C and ^{15}N in
85 biological tissues across trophic levels is variable depending on the species involved,
86 commonly used trophic enrichment factors (TEFs) (ca. 1‰ for ^{13}C and 3 to 4‰ for ^{15}N ; e.g.
87 DeNiro & Epstein 1978, Minagawa & Wada 1984) in place of species-specific ones (e.g.
88 Yokoyama et al. 2005) may mislead the interpretation of food-chain structure. In addition to
89 bulk analysis, $\delta^{15}\text{N}$ of amino acids (AAs) have been widely used to estimate more accurate TPs
90 (McClelland & Montoya 2002, Ohkouchi et al. 2017 and references therein). Among the AAs
91 found in organisms, 2 types have been identified in terms of their ^{15}N -enrichment across
92 trophic levels. One is ‘trophic amino acids’, the $\delta^{15}\text{N}$ of which significantly increase from one
93 trophic level to the next, while the other, ‘source amino acids’, show only slight shifts in $\delta^{15}\text{N}$
94 between trophic levels. Therefore, the analysis of $\delta^{15}\text{N}$ in a set of AAs covering these 2 types
95 (mainly phenylalanine for source AA and glutamic acid for trophic AA) for a target species
96 potentially enables the identification of accurate TPs (Chikaraishi et al. 2009). Furthermore, it
97 has recently been reported that one trophic AA, alanine, is effective at more correctly
98 estimating a trophic pathway including both protistan and metazoan consumers
99 (Gutiérrez-Rodríguez et al. 2014, Décima et al. 2017, Landry & Décima 2017). Therefore, the

100 use of alanine as trophic AA would be more adequate for food chain analyses in coastal oceans
101 as well as pelagic waters where trophic transfers through protistan grazers are not negligible.

102 Callianassid shrimp or ghost shrimp (Pleocyemata: Axiidea) play key roles as
103 ecosystem engineers and community organizers in estuarine and coastal marine sedimentary
104 habitats (Pillay & Branch 2011, Dworschak et al. 2012). Planktotrophic larvae of callianassids
105 go through 3 to 7 zoeal stages to reach the decapodid stage (Kubo et al. 2006, Pohle et al. 2011,
106 Kornienko et al. 2015). The survival rate of those larvae significantly affects the population
107 dynamics of adults, which could further exert an impact on the community and ecosystem
108 surrounding that population. Following the release of zoeae I from the shore, those larvae grow
109 in the inner-shelf coastal ocean (Johnson & Gonor 1982, Yannicelli et al. 2006, Tamaki et al.
110 2010).

111 In midwestern Kyushu, Japan, a population of the callianassid shrimp *Nihonotrypaea*
112 *harmandi* (Bouvier 1901) is distributed on intertidal sandflats in a region ranging from estuary
113 (Ariake Sound) to inner-shelf coastal ocean (Amakusa-nada) (Tamaki et al. 1999) (Fig. 1).
114 Around the western edge of the region, the largest local population in the region exists on the
115 intertidal sandflat in Tomioka Bay (Tomioka sandflat; 32.521° N, 130.037° E) (Tamaki &
116 Harada 2005, Tamaki & Takeuchi 2016). Tomioka Bay forms a coastal boundary layer with
117 water depths ≤ 30 m. Larvae released from the sandflat are transported across Tomioka Bay to
118 Amakusa-nada, where the main nursery ground for larvae lies 10 to 20 km north to west of the
119 sandflat (Tamaki & Miyabe 2000, Tamaki et al. 2010, 2013). Complete rearing of *N. harmandi*
120 larvae has been achieved in the laboratory using the rotifer *Brachionus rotundiformis*, and
121 newly hatched *Artemia* sp. nauplius (Konishi et al. 1999). By contrast, examination of both the
122 mouthpart morphology and digestive tract content of field-caught *N. harmandi* larvae suggests
123 that they are herbivorous, mainly feeding on diatoms (Somiya et al. 2014). But there are no
124 records of complete larval rearing solely with phytoplankton.

149 of 21 to 22°C and salinities of 33.5 to 34 during the end of July to early August 2006 (Tamaki
150 et al. 2010). In the laboratory at 21°C, it took 30 d for most of the newly hatched zoeae reared
151 with *Brachionus rotundiformis*, and newly hatched *Artemia* sp. nauplii to reach the decapodid
152 stage (Tamaki et al. 2013). Starved zoeae I survived for up to 5 d and never proceeded to the
153 zoea II stage (Y. Saitoh & A. Tamaki unpubl. data).

154 **Setup for ghost shrimp larval rearing in the laboratory**

155 Larvae of *N. harmandi* were mass-reared in 2 black, semi-cylindrical polyethylene
156 tanks (rearing tanks 1 and 2) with a 30 l capacity during July and August 2013. Surface water
157 from a coastal ocean area, filtered through a 10 µm mesh (TOCEL; JFE Advantech), was
158 stored and used for the rearing experiment. The water temperature in the tanks was kept at
159 around 22°C (which corresponded to the field water-column conditions for the larvae) using 2
160 water baths (see Section 1.1 in the Supplement at
161 www.int-res.com/articles/suppl/mXXXXpXXX_supp.pdf). The seawater temperature and
162 salinity in each tank were recorded every 1 min with a Compact-CT (JFE Advantech). Gentle
163 aeration and slow water circulation in each tank was made with a stream of tiny air bubbles
164 passed through 2 air stones with 20 mm ϕ , which kept only food particles (i.e. not deposits)
165 suspended. During rearing, the room lights were kept off as a rule, except during water
166 exchanges, food supply, and retrieval of larvae. Furthermore, the tank tops were covered with
167 fine-mesh black sheets, which was intended to approximate faint daylight conditions at 30 to
168 40 m water depths in Amakusa-nada (Tamaki et al. 2010).

169 **Collection of ghost shrimp ovigerous females and initiation of larval rearing**

170 During daytime low tide on the Tomioka sandflat on 7 July 2013, 57 females of *N.*
171 *harmandi* with well-developed embryos were collected. These embryos were expected to hatch
172 during the immediate nighttime as determined by their large eyes and transparent bodies due to

173 a minimum amount of remaining yolks. In the laboratory, the females were kept in containers
174 with filtered seawater until their larvae were released. Synchronized mass-release events
175 occurred twice, at 02:20 and 04:00 h on 8 July. Each time, swimming larvae (newly hatched
176 zoeae I) were collected with a nylon net and gently transferred to a transparent container
177 containing 1.5 to 3.0 l filtered seawater. From each container, 12 samples of 5 ml water were
178 randomly taken with a measuring pipette. After counting the number of larvae in each sample,
179 they were returned to the container for re-sampling. The mean (\pm SD) numbers of larvae
180 sample⁻¹ (12.0 ± 4.0 and 3.7 ± 1.2 in the first and second events, respectively) were used to
181 estimate the total numbers of larvae, which were approximately 7200 and 1100 larvae in the
182 first and second events, respectively. These 2 batches of larvae were mixed in a cup with
183 filtered seawater and split into 2 cups each with a known volume of water. It was estimated that
184 4500 and 3800 larvae were contained in these cups. In this order, the cup contents were gently
185 poured into rearing tanks 1 and 2, respectively, each of which contained a sufficient quantity of
186 pure-cultured diatoms (*C. gracilis*; Bivalve Culture Institute, Sasebo city). All the above
187 procedures were completed by 05:00 h. The live diatom stock (approximately 5×10^8 cells ml⁻¹
188 in a bottle of 1000 ml seawater) can be stored at between 0 and 5°C in the refrigerator for at
189 least 50 d (see Yokoyama et al. 2005 and Tamaki et al. 2013 for successfully rearing
190 decapodids and juveniles of *N. harmandi* with this product).

191 **Routine procedures for ghost shrimp larval rearing and retrieval, and data** 192 **analysis**

193 Starting from 09:00 h on 8 July 2013 (Day 0), the daily routine work for each of the 2
194 tanks was completed between 09:00 and 11:00 h. Carcasses, feces, and exuviae of *N. harmandi*
195 larvae, and *C. gracilis* flocs deposited on the tank bottom were removed with pipettes.
196 Three-fourths of the tank water was gently siphoned off, with a 63 μ m mesh nylon net covering
197 the siphon entrance to prevent larvae from being inhaled, and replaced with new filtered

222 The multiple cohorts of decapodids of *N. harmandi* occurring through the rearing dates
223 might reflect those cohorts that could exist already at the stages of newly fertilized embryos
224 and newly hatched zoeae I, with dimension differences between the cohorts. The subsequent
225 developmental durations can vary accordingly. To detect the presence of such initial cohorts,
226 females with (1) newly fertilized embryos and (2) embryos about to hatch were collected from
227 the Tomioka sandflat during daytime low tides on 24 and 26 August and 10 September 2014.
228 Determination of these embryonic states is detailed in Section 1.2 in the Supplement. The
229 females with embryos-(1) were individually fixed with 10% neutralized seawater formalin.
230 Those with embryos-(2) were brought alive to the laboratory.

231 Under a light microscope in the laboratory, the above embryonic-state-(1) was
232 confirmed as the stage before reaching the embryonic nauplius. In total, 73 females with those
233 embryos were obtained. Ten embryos were randomly removed from each female, and the
234 longer and shorter axes of the ellipse of each embryo in plane aspect were measured to 0.01
235 mm with an eyepiece micrometer attached to the microscope at 100× magnification. The mean
236 longer and shorter axis lengths were used to calculate the volume of a spheroid embryo with its
237 longer axis as the axis of revolution. Using data for all females combined, the embryonic
238 volume-frequency distribution was made. The females with embryonic-state-(2) were kept
239 individually in small containers with filtered seawater until their larval release during the
240 immediate nighttime. In total, 61 females released larvae, which were collected and fixed with
241 5% neutralized seawater formalin. For 10 randomly chosen larvae from each female, the
242 mid-dorsal total length in lateral aspect, from rostral spine to telson tips, was traced with a
243 camera lucida of a stereomicroscope at 30 to 50× magnification. The curve length was
244 measured to 0.01 mm using ImageJ 1.48v (<http://imagej.nih.gov/ij/>). Using the mean larval
245 total length from every female, a total-length-frequency distribution was made. Each
246 composite normal-distribution group was separated into the components as previously.

247 **Sampling diatoms and larvae reared in the laboratory for SI analysis**

248 The liquid with concentrated diatoms (*C. gracilis*) for the rearing of *N. harmandi* larvae
249 was sampled for SI analysis on Days 0, 9, 15, and 28, and immediately processed (see Section
250 1.3 in the Supplement).

251 The numbers of *N. harmandi* larvae retrieved for SI analysis (combined from the 2
252 rearing tanks) consisted of approximately 50 to 60 for zoeae I immediately after hatching
253 (retrieved separately from those larvae subsequently used for the rearing), 9 for zoeae IV on
254 Day 15, 52 for decapodids (7 to 16 each on Days 28 and 30 to 33), and all zoeae remaining on
255 Day 40 (20 zoeae V and 37 zoeae VI). After their defecation, these specimens were kept frozen
256 at -20°C until analysis. The TEF associated with the larval feeding on diatoms was calculated
257 as the difference in SI values between zoeal whole-body tissues and diatoms as follows, when
258 the larval SI values became steady during the rearing: $\text{TEF } (\Delta\delta^{13}\text{C or } \Delta\delta^{15}\text{N}) = \text{SI value}$
259 $(\text{larvae}) - \text{SI value} (\text{diatoms})$.

260 **Field sampling for SI and plankton composition analyses**

261 Water and biological sampling for chemical analyses were carried out at Stn A in
262 Amakusa-nada (Fig. 1) during 2 cruises in 2012 (10 to 12 July and 7 to 10 August) and 3
263 cruises in 2013 (27 to 29 June, 10 to 12 July, and 7 to 9 August) onboard the training vessel
264 (T/V) 'Kakuyo-Maru' (Nagasaki University) equipped with 12 Niskin bottles (5 l) mounted on
265 a rosette multiple sampler (RMS; General Oceanics) with the CTD probe (SBE 9/11; Seabird
266 Electronics). Regarding the potential food sources for *N. harmandi* larvae, water samples for
267 particulate organic matter (POM) to be used for SI analyses were collected, 2 or 3 times in each
268 cruise, from the chlorophyll (chl) maximum layers and other 1 or 2 layers 5 to 10 m above or
269 below each chl maximum layer (10 to 25 m depth except for August 2012 [30 to 50 m], as
270 detected with the CTD probe), which was expected to contain high content of fresh

271 phytoplankton. Chl *a* concentrations in the water column ($\mu\text{g l}^{-1}$) were calculated from chl
272 fluorescence at each layer, and conversion factor was obtained at Stn A during the shared
273 cruises in August and September 2012 (S. Takeda pers. comm.). Each 5 to 10 l sample of
274 seawater was filtered through a 200 μm nylon mesh and a pre-combusted filter (GF/F; 47 mm
275 ϕ), and then the sample was gently washed with a few ml of distilled water. Samples were kept
276 frozen at -20°C until analysis and processed in the same manner as for the diatom sample (see
277 Section 1.3 in the Supplement).

278 Zoeae of *N. harmandi* (no decapodids were obtained) and their potential food sources,
279 zooplankton (mostly composed of calanoid copepods), were collected for bulk SI analyses by
280 vertical towing with a Norpac Net (0.45 m mouth diameter, 1.8 m length, and 330 μm mesh
281 size) from 5 m above the seabed to the surface at Stn A at both low- and high-tide times. These
282 plankton samples were immediately fixed with neutralized formalin solution (5% final conc.)
283 and brought to the laboratory. Under a stereomicroscope, larval samples were separated into
284 the 6 stages (zoea I to VI) both in 2012 and 2013, while zooplankton were collected only in
285 2012 and separated into 6 body-size fractions (3.5–2.5, 2.5–2.0, 2.0–1.5, 1.5–1.2, 1.2–0.8, and
286 0.8–0.3 mm). Prior to the SI analysis of these field-collected samples, the effect of preservation
287 in formalin solution on the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values for zoeal tissues (i.e. difference in $\delta^{13}\text{C}$ and
288 $\delta^{15}\text{N}$ values between the zoeal samples with and without formalin treatment) was tested for
289 several zoeal stages and the values used for the calibration of the SI values for both zoeae and
290 other zooplankton.

291 Zoeae of *N. harmandi* for $\delta^{15}\text{N}$ analysis in amino acids were collected on 10 August
292 2012 from 30 m depth by horizontally towing a ‘fish-larvae’ net (1.3 m mouth diameter, 4.5 m
293 length, and 330 μm mesh size) at a speed of 1.5 knots for 10 min. The sample was immediately
294 brought to the laboratory alive, and zoeae VI were selected under a stereomicroscope and kept
295 frozen at -20°C until analysis. About a 12 h lag from sampling to freezing was enough for

296 those zoeae to evacuate their digestive tract contents, suggesting no contamination of zoeal
297 tissues with diet/gut contents.

298 Water samples for the taxonomic composition analysis of nano- to micro-sized
299 plankton were collected from 4 depth layers (0, 20, 40, and 60 m) with a bucket (0 m) or Niskin
300 bottles on 7 August 2012. Sampled water was immediately fixed with acid Lugol's solution
301 (2% in final conc.) and stored in cool and dark conditions. In the laboratory, 3.1 to 4.4 ml of
302 water sample was set in a Sedgewick–Rafter chamber (Guillard 1978) for each layer, so that
303 >300 of those plankton were included. All individuals larger than 4 μm in equivalent spherical
304 diameter (i.e. eukaryotic unicellular organisms and filamentous cyanobacteria) were observed
305 and identified to varying levels from class to suborder under an inverted biological microscope
306 equipped with a 60 \times objective lens (Olympus IX71). Plankton biovolume was individually
307 estimated assuming approximate geometrical figures such as spherical, ellipsoidal, cylindrical,
308 and conical shapes.

309 **POM degradation experiment**

310 To check for the potential alteration of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values in POM by microbial
311 degradation in the water column during suspension and sinking into deeper layers, laboratory
312 incubation experiments were conducted in June and July 2013. To prepare POM presumably
313 with a high content of phytoplankton, seawater samples collected from the chl maximum layer
314 at Stn A (preceding section) were filtered sequentially through 200 and 20 μm nylon meshes.
315 The organic matter trapped on the 20 μm nylon mesh was re-suspended in filtered seawater
316 (through GF/F) in duplicate 200 ml bottles. The sample bottles were immediately stored in a
317 portable incubator (CN-25C; Mitsubishi Electric) onboard and incubated at a temperature of
318 22 $^{\circ}\text{C}$ in the dark for 1 wk. Duplicate POM samples were collected on pre-combusted GF/F
319 filters (25 mm ϕ) on 3 dates (Days 0, 2 or 3, and 5 or 7), and processed in the same manner as
320 for the diatom samples in the larval rearing (see Section 1.3 in the Supplement).

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AA extraction and purification

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Roughly powdered zoecal whole-body tissues of *N. harmandi* (zoea VI, 2 sets of about 20 ind., ca. 4 mg dry weight collected at Stn A in August 2012) were pre-treated prior to $\delta^{15}\text{N}$ analysis of individual AAs. The subsequent procedures followed the protocol given in Ishikawa et al. (2014), with some modifications (see Section 1.4 in the Supplement).

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SI measurement of bulk and AA samples

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For the zooplankton samples (including *N. harmandi* larvae) collected at Stn A in Amakusa-nada in 2012 and 2013, and from the rearing experiment for the larvae, approximately 0.3 to 0.5 mg of the sample in dry weight (i.e. 15 to 20 ind. for zoea I, 11 to 13 for zoea II, 9 to 11 for zoea III, 6 to 8 for zoea IV, 3 to 5 for zoea V, 3 each for zoea VI and decapodid, and 10 to 50 ind. for zooplankton depending on body-size groups, randomly selected from the sample stock) was transferred to silver capsules (SÄNTIS Analytical) and dried at 60°C after acidification with a few drops of 1 N HCl to remove inorganic carbon. The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values for diatoms (diet in the rearing experiment), POM (field sample and degradation experiment), and zooplankton samples were measured with an elemental analyzer and an isotope-ratio mass spectrometer (FLASH 2000-Conflo IV-Delta V Advantage; Thermo Fisher Scientific). Instrument precision was checked with a calibration standard (L-alanine) every 5 samples (standard deviation <0.12‰ for $\delta^{13}\text{C}$, <0.14‰ for $\delta^{15}\text{N}$). The $\delta^{15}\text{N}$ values for the samples with lower N contents (<20 μg) were eliminated from the results (e.g. N at POM degradation experiment), because the accuracy of the measured values can be low, despite the calibration with standards, due to a mass-dependent shift in $\delta^{15}\text{N}$ values in small-quantity samples (Ogawa et al. 2010).

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The $\delta^{15}\text{N}$ values of the 3 kinds of AAs (alanine [Ala], glutamic acid [Glu], and phenylalanine [Phe]) were determined by gas chromatography/combustion/isotope ratio mass

345 spectrometry (GC/C/IRMS) using a Delta V plus isotope ratio mass spectrometer (Thermo
346 Fisher Scientific) coupled to a gas chromatograph (GC7890A; Agilent Technologies) via a
347 modified GC-Isolink interface consisting of combustion and reduction furnaces. The AA
348 derivatives were injected into the GC column using a Gerstel PTV injector in solvent vent
349 mode. The programs on temperature, retention time, and carrier gas flow rate in each process
350 (i.e. injection, combustion, reduction, and separation in GC) followed the methods given in
351 Ishikawa et al. (2014). The CO₂ generated in the combustion furnace was removed using a
352 liquid N trap. Standard mixtures of 8 kinds of AAs ($\delta^{15}\text{N}$ ranging from -25.9 to $+45.6\text{‰}$) were
353 analyzed with zoal sample in turn to confirm the reproducibility of the isotope measurements.
354 Standard deviations of the standards were better than 0.8‰ with a sample quantity of 5 nmol
355 μl^{-1} N.

356 It was anticipated that protozoan consumers potentially contributed to the diet of *N.*
357 *harmandi* larvae (cf. Fileman et al. 2014); there is a certain amount of biomass of protozoa such
358 as ciliate and heterotrophic dinoflagellate in coastal waters, and sinking phytodetritus with
359 heterotrophs could be a candidate for the diet. Therefore, both the TP-estimate based on the
360 $\delta^{15}\text{N}$ values of Glu and Phe with canonical parameters for the metazoan food chain (i.e.
361 $\text{TP}_{\text{Glu/Phe}}$; Chikaraishi et al. 2009) and the TP-estimate based on the $\delta^{15}\text{N}$ values of Ala and Phe
362 with another set of parameters for the metazoan food chain potentially including a protozoan
363 pathway (i.e. $\text{TP}_{\text{Ala/Phe}}$; Décima et al. 2017) were calculated for the targeted samples using the
364 following equations, respectively:

$$365 \quad \text{TP}_{\text{Glu/Phe}} = (\delta^{15}\text{N}_{\text{Glu}} - \delta^{15}\text{N}_{\text{Phe}} - 3.4) / 7.6 + 1 \quad (1)$$

$$366 \quad \text{TP}_{\text{Ala/Phe}} = (\delta^{15}\text{N}_{\text{Ala}} - \delta^{15}\text{N}_{\text{Phe}} - 3.2) / 5.5 + 1 \quad (2)$$

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RESULTS

369 **Size-frequency distributions for ghost shrimp embryos and newly hatched**
370 **zoeae I**

371 The newly fertilized embryo-volume-frequency distribution and newly hatched zoea
372 I-total length-frequency distribution of *Nihonotrypaea harmandi* in 2014 are shown in Fig. 2.
373 Each frequency distribution comprised 2 cohorts (cohorts 1 and 2), to which normal
374 distributions were fitted. The mean diameters of embryos (per female) ranged from 0.44 (short
375 axis) to 0.52 mm (long axis). The estimated embryo volume ranged from 0.37 to 0.59 mm³.
376 Cohort 1 comprised 77% of all embryos, with mean (\pm SD) of 0.42 ± 0.02 mm³. For cohort 2,
377 the mean embryo volume was 0.50 ± 0.04 mm³, which was 1.19 times greater than for cohort 1.
378 The total length of zoea I ranged from 2.56 to 2.95 mm. Cohort 1 comprised 54% of all zoeae I,
379 with mean of 2.70 ± 0.06 mm. For cohort 2, the mean total length was 2.87 ± 0.06 mm. The
380 (mean total length)³ was 1.20 times greater than that in cohort 1, where the cube is a measure
381 for larval volume.

382 **Water temperature and salinity in laboratory rearing tanks for ghost**
383 **shrimp larvae**

384 The time series of seawater temperature and salinity in the 2 rearing tanks for *N.*
385 *harmandi* larvae through the whole period (05:00 h on 8 July to 09:00 h on 17 August in 2013)
386 are shown in Fig. S1 in the Supplement. In rearing tank 1, the water temperature ranged from
387 20.9 to 23.1°C, with mean (\pm SD) of 22.2 ± 0.25 °C. Salinity ranged from 32.5 to 34.1, with
388 mean of 33.5 ± 0.2 . In rearing tank 2, the water temperature ranged from 21.3 to 23.0°C, with
389 mean of 22.3 ± 0.1 °C. Salinity ranged from 32.3 to 33.8, with mean of 33.4 ± 0.2 .

390 **Development and survival of ghost shrimp larvae in laboratory rearing**

391 At the time of daily exchange of water in the rearing tanks, the ‘old’ water was
392 brownish in color, suggesting an excess ration of *Chaetoceros gracilis*. The daily occurrence of

393 decapodids of *N. harmandi* in rearing tanks 1 and 2 is shown in Fig. 3. During the rearing
394 period in each tank, there were 2 major cohorts (cohorts 1 and 2), to which normal distribution
395 curves were fitted. Decapodids emerged first on Day 26 in both tanks. In rearing tank 1, a total
396 of 175 decapodids occurred until Day 40, giving a survival rate of 3.9% of the initial number of
397 zoeae I from which the number of advanced-stage zoeae retrieved on the way to the decapodid
398 stage was subtracted. Cohort 1 comprised 30% of all decapodids, with mean occurrence in day
399 number being 28.7 ± 1.0 . For cohort 2, the mean (\pm SD) occurrence as day number was $34.2 \pm$
400 2.6 . In rearing tank 2, a total of 127 decapodids occurred until Day 40, giving a survival rate of
401 3.4%. Cohort 1 comprised 35% of all decapodids, with mean occurrence in day number being
402 28.8 ± 1.4 . For cohort 2, the mean occurrence as day number was 35.3 ± 2.3 .

403 **TEF associated with diatom-feeding by ghost shrimp larvae**

404 The SI values of the diet (diatom: *C. gracilis*) in the *N. harmandi* larval rearing
405 experiment were almost constant through the study period (mean \pm SD, $n = 17$ values of $\delta^{13}\text{C} =$
406 $-22.7 \pm 0.3\text{‰}$ and $\delta^{15}\text{N} = -1.9 \pm 0.3\text{‰}$; Fig. 4), and these mean values were used for the
407 calculation of TEFs. The SI values in the zoeal whole-body tissues gradually shifted, with
408 development, from those in zoea I ($\delta^{13}\text{C} = -18.1 \pm 0.1\text{‰}$ and $\delta^{15}\text{N} = 7.1 \pm 0.02\text{‰}$, $n = 3$) to the
409 asymptotic values closer to that for diatoms. The SI values in the body tissue were -20.7‰ for
410 $\delta^{13}\text{C}$ and 1.0‰ for $\delta^{15}\text{N}$ ($n = 1$) at the zoea IV on Day 15 and reached a steady state at the
411 decapodid during Day 28 to Day 33 ($\delta^{13}\text{C} = -20.7 \pm 0.3\text{‰}$ and $\delta^{15}\text{N} = 0.0 \pm 0.3\text{‰}$, $n = 15$). The
412 SI values for the larvae that still remained at the zoeal stages (V and VI) in the tanks on Day 40
413 ($\delta^{13}\text{C} = -20.5 \pm 0.3\text{‰}$ and $\delta^{15}\text{N} = 0.0 \pm 0.3\text{‰}$, $n = 3$) were similar to those steady values. Thus,
414 the TEFs associated with the feeding of diatoms by *N. harmandi* zoeae was calculated as 2.0‰
415 for $\delta^{13}\text{C}$ and 1.9‰ for $\delta^{15}\text{N}$.

416 **Measurement for shift in phytoplankton SI values with degradation**

417 During the degradation experiment in June 2013, the mean SI values for the POM
418 (fraction between 20 and 200 μm) from the chl maximum layers gradually decreased from Day
419 0 (mean [\pm SD] $\delta^{13}\text{C} = -20.0 \pm 0.4\text{‰}$, $n = 3$ and $\delta^{15}\text{N} = 5.6$ to 5.7‰ , $n = 2$) to Day 7 ($\delta^{13}\text{C} =$
420 -20.9 to -20.3‰ , $n = 2$ and $\delta^{15}\text{N} = 4.9\text{‰}$, $n = 1$) (Fig. 5). A similar decreasing tendency was
421 observed in July 2013. The SI values for the POM decreased from Day 0 ($\delta^{13}\text{C} = -19.4$ to
422 -19.1‰ , $n = 2$ and $\delta^{15}\text{N} = 6.2$ to 6.4‰ , $n = 2$) to Day 5 ($\delta^{13}\text{C} = -19.6$ to -19.5‰ , $n = 2$ and
423 $\delta^{15}\text{N} = 5.2$ to 5.6‰ , $n = 2$). The average daily decreasing rates (‰ d^{-1}) were -0.09 for $\delta^{13}\text{C}$ and
424 -0.11 for $\delta^{15}\text{N}$ during the 7 d in June, and -0.07 for $\delta^{13}\text{C}$ and -0.18 for $\delta^{15}\text{N}$ during the 5 d in
425 July.

426 **Estimation of TP for ghost shrimp larvae based on AA $\delta^{15}\text{N}$**

427 The AA-specific $\delta^{15}\text{N}$ values of 2 sets of the *N. harmandi* zoea-VI samples were 2.3
428 and 2.9 ‰ for Phe, 14.7 and 15.2 ‰ for Ala, and 14.4 and 14.5 ‰ for Glu, respectively. The
429 calculated mean TPs for ghost shrimp larvae were 2.1 for $\text{TP}_{\text{Glu/Phe}}$ and 2.7 for $\text{TP}_{\text{Ala/Phe}}$.

430 **Taxonomic composition and standing crop of small-sized plankton in** 431 **Amakusa-nada**

432 Total abundance and total biovolume of nano- to micro-sized plankton in the 4 depth
433 layers from the surface to 60 m at Stn A on 7 August 2012 ranged from 7.0×10^4 to 1.2×10^5
434 ind. l^{-1} and from 1.5×10^8 to $2.4 \times 10^8 \mu\text{m}^3 \text{ l}^{-1}$, respectively, with values tending to decrease
435 with depth (Fig. 6). Diatoms were dominant through the water column, both in abundance
436 (47.3 to 55.3%) and biovolume (37.7 to 80.7%). Specifically, centric diatoms composed of
437 Coscinodiscineae, Rhizosoleniineae, and Biddulphiineae accounted for 70.7 to 90.0% in
438 abundance and 93.6 to 97.2% in biovolume of the diatoms. Dinoflagellates (11.0 to 27.7% in
439 abundance and 9.5 to 23.1% in biovolume, mainly composed of Gymnodiniales) or ciliates (5.8

440 to 12.2% in abundance and 3.8 to 33.2% in biovolume, mainly composed of aloricate type)
441 were subordinate.

442 **Bulk SI values for ghost shrimp larvae and their potential food sources**

443 The salinity averaged over the surface to 40 m depth at Stn A (Fig. 1) from CTD data in
444 each cruise was lower in 2012 (mean [\pm SD]: 33.0 ± 0.5 in July and 32.6 ± 0.3 in August) than in
445 2013 (33.7 ± 0.2 in June, 33.6 ± 0.2 in July, and 33.4 ± 0.1 in August). Irrespective of the years,
446 around the chl maximum layers at Stn A, higher chl *a* concentrations ($>10 \mu\text{g l}^{-1}$) were
447 observed at 10 to 20 m depths in June and July, while lower peaks of 2 to $8 \mu\text{g l}^{-1}$ were
448 observed at 30 to 50 m depths in August.

449 The SI values of biological samples and POM from Amakusa-nada were compiled
450 separately for each of 2012 and 2013 (Fig. 7). The $\delta^{13}\text{C}$ in *N. harmandi* zoeal whole-body
451 tissues preserved in formalin ($\delta^{13}\text{C}_{\text{formalin}}$) had lighter values than those in frozen (= 'intact')
452 samples ($\delta^{13}\text{C}_{\text{frozen}}$) (i.e. mean [\pm SD] $\Delta\delta^{13}\text{C}$ [$\equiv \delta^{13}\text{C}_{\text{formalin}} - \delta^{13}\text{C}_{\text{frozen}}$] = $-1.0 \pm 0.2\text{‰}$, $n = 9$,
453 while $\delta^{15}\text{N}_{\text{formalin}}$ had heavier values than those in $\delta^{15}\text{N}_{\text{frozen}}$ i.e. $\Delta\delta^{15}\text{N}$ [$\equiv \delta^{15}\text{N}_{\text{formalin}} - \delta^{15}\text{N}_{\text{frozen}}$]
454 = $0.5 \pm 0.3\text{‰}$, $n = 9$). Therefore, the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values for the zoeal whole-body tissues
455 given in the food-chain diagram (Fig. 7) are those after the correction for the effect of formalin
456 preservation by subtracting the above mean Δ -values from $\delta^{13}\text{C}_{\text{formalin}}$ or $\delta^{15}\text{N}_{\text{formalin}}$. The $\delta^{13}\text{C}$
457 and $\delta^{15}\text{N}$ values for other zooplankton whole-body tissues also designate those compensated
458 ones.

459 In 2012, the mean $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values in *N. harmandi* zoeal whole-body tissues at
460 stages I and II inclusive were $-18.3 \pm 0.2\text{‰}$ and $6.0 \pm 0.2\text{‰}$ ($n = 7$), respectively, and those
461 values at stages III and IV inclusive had similar values ($\delta^{13}\text{C}$: $-18.5 \pm 0.3\text{‰}$; $\delta^{15}\text{N}$: $6.3 \pm 0.2\text{‰}$,
462 $n = 6$) (Fig. 7a). Most of the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values in the zoeae at stages V and VI inclusive
463 showed lighter values (i.e. $\delta^{13}\text{C}$: $-20.2 \pm 0.3\text{‰}$; $\delta^{15}\text{N}$: $5.7 \pm 0.2\text{‰}$, $n = 5$), which were

464 sufficiently distant from those at the earlier stages, though several latest-stage specimens (2 ind.
465 of zoea V and 1 ind. of zoea VI) took values closer to the earlier-stage values. The $\delta^{13}\text{C}$ and
466 $\delta^{15}\text{N}$ values for POM around the chl maximum layer showed large variations ranging from
467 -22.8 to -18.7‰ ($-20.5 \pm 1.3\text{‰}$, $n = 10$), and from 4.8 to 6.2‰ ($5.6 \pm 0.4\text{‰}$, $n = 10$),
468 respectively. The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values in the smaller zooplankton (0.3 to 1.2 mm in size) were
469 $-19.9 \pm 0.3\text{‰}$ ($n = 3$) and $5.5 \pm 0.2\text{‰}$ ($n = 3$), respectively, while the larger zooplankton (1.2 to
470 3.5 mm in size) had relatively heavier values ($-19.5 \pm 0.2\text{‰}$, $n = 8$ for $\delta^{13}\text{C}$ and $6.3 \pm 0.3\text{‰}$, n
471 $= 8$ for $\delta^{15}\text{N}$).

472 In 2013, the mean $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values in *N. harmandi* zoeal whole-body tissues at
473 stages I and II inclusive were -17.6 and 6.1‰ ($n = 1$ for June and July each), respectively, and
474 those values gradually shifted to lighter ones with larval development; at stages III and IV
475 inclusive ($\delta^{13}\text{C}$: $-18.3 \pm 0.3\text{‰}$; $\delta^{15}\text{N}$: $6.0 \pm 0.2\text{‰}$, $n = 6$) and at stages V and VI inclusive ($\delta^{13}\text{C}$:
476 $-18.7 \pm 0.2\text{‰}$; $\delta^{15}\text{N}$: $5.7 \pm 0.3\text{‰}$, $n = 10$) (Fig. 7b). As in 2012, the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values for
477 POM in the chl maximum layer in 2013 showed large variations ranging from -22.1 to -19.4‰
478 ($-20.5 \pm 1.0\text{‰}$, $n = 5$), and from 4.1 to 5.6‰ ($5.0 \pm 0.6\text{‰}$, $n = 5$), respectively. The values for
479 POM in 2012 and 2013 inclusive were near to the previously reported values at 5 m depth on a
480 location 7.5 km east of Stn A in May and June 2004 ($\delta^{13}\text{C} = -21.3 \pm 0.2\text{‰}$ and $\delta^{15}\text{N} = 5.9 \pm$
481 0.3‰ ; Shimoda et al. 2007).

482 DISCUSSION

483 The present study was initiated primarily to fill a gap between the universally standard
484 laboratory diet (i.e. excess ration of a specific set of microzooplankton) for rearing decapod
485 crustacean larvae and the absence, or low availability, of nutritionally equivalent
486 microzooplankton in the coastal ocean water column containing plentiful phytoplankton and
487 their detritus (phytodetritus).

488 The larvae of *Nihonotrypaea harmandi* mass-reared at a water temperature of 22°C,
489 with a diet consisting solely of pure-cultured nano-sized diatoms *Chaetoceros gracilis* in an
490 excess ration, reached the first peak in decapodid occurrence on Day 29 and the second one on
491 Days 34 and 35, and survived at rates of 3.4 to 3.9% from the initially housed zoeae I (Fig. 3).
492 Zoeal bulk SI equilibration with regard to diatom SIs (Fig. 4) excludes the possibility of
493 ingesting deposited animal material. The above-mentioned values are comparable with those
494 obtained from the 2 previous rearing experiments under similar conditions to the present study
495 (Tamaki et al. 2013). There, about 6600 newly hatched zoeae were housed in a 30 l tank and
496 reared with a combination of *C. gracilis* (targeted only for zoea I), *Brachionus rotundiformis*
497 (rotifer; provided for all zoeal stages), and *Artemia* sp. nauplii (targeted for zoeae III-VI), with
498 each food item in an excess ration at 2 constant water temperatures. At 21°C, the first and
499 second peaks in decapodid occurred on Day 30 (Tamaki et al. 2013) and Day 35 (A. Tamaki
500 unpubl. data), respectively, with a combined retrieval rate of 7.35% (Tamaki et al. 2013). At
501 24°C, those peaks occurred on Day 30 (Tamaki et al. 2013) and Day 34 (A. Tamaki unpubl.
502 data), with a combined retrieval rate of 4.7% (Tamaki et al. 2013). The 2 cohorts in the *N.*
503 *harmandi* decapodids are likely to have originated from those cohorts in newly fertilized
504 embryos and newly hatched zoeae I, for which a common ratio in the larger to smaller mean
505 volumes existed (i.e. 1.2; Fig. 2). The larger eggs in *N. harmandi* occur around the time when
506 females begin to participate in reproduction, and repeated egg laying with time brings about
507 smaller eggs (A. Tamaki unpubl. data; cf. Kubo et al. 2006). One congeneric species, *N.*
508 *japonica*, has a mean egg volume 1.5 to 2.0 times greater and a mean zoea I total length 1.11
509 times greater than those in *N. harmandi* (see Kubo et al. 2006 and Tamaki & Miyabe 2000,
510 respectively); the difference in cubic total length is 1.37 times. The larval development of *N.*
511 *japonica*, from zoea I through zoea II to zoea V to the first occurrence of decapodids, took 16 d
512 in a laboratory rearing under similar conditions as in Tamaki et al. (2013) (Miyabe et al. 1998).
513 Among decapod crustacean species, those with larger eggs tend to have greater yolk contents

514 (Wear 1974), which would lead to shorter larval developmental durations, including
515 callianassid shrimp species (Kubo et al. 2006). This might also be true for the intraspecific
516 difference in egg volume and larval developmental duration.

517 Of the 3 available studies about the successful rearing of larvae of the decapod
518 crustacean suborder Pleocyemata solely with phytoplankton, 2 were qualitative, in which
519 nano- to micro-sized diatoms and dinoflagellates (Atkins 1955) or green alga and diatom
520 (Bousquette 1980) were provided to pinnotherid brachyurans. The other one quantitatively
521 tested the effect of 3 diatom species with different sizes (250, 40, and 10 μm in ϕ) on the
522 brachyuran *Hyas coarctatus*, with only the largest species (*Biddulphia sinensis*: micro-size)
523 found to be a valid food but much less efficient than *Artemia* sp. nauplii (Harms & Seeger
524 1989). In that case, chain-forming in diatoms was regarded as essential for ingestion by the
525 zoeae. In the digestive tract contents of *N. harmandi* zoeae from Amakusa-nada, micro-sized
526 diatoms belonging to Rhizosoleniineae, Coscinodiscineae, Biddulphiineae, and Pennales were
527 found (Somiya et al. 2014). Each diatom frustule or piece was shorter than 25 μm in width and
528 height, while the length was not limited; smaller or slender diatoms might be ingested whole,
529 while larger or wider diatoms would be masticated by the mandible before ingestion. Cells of *C.*
530 *gracilis* used for the present study were nano-sized and not chain-forming, and *N. harmandi*
531 zoeae ingest them not individually but as a bolus accumulated by filter-feeding around the
532 mouthparts (R. Somiya pers. comm.). Such ingestion of lumps was observed for conspecific
533 decapodids and juveniles (Yokoyama et al. 2005, Tamaki et al. 2013). Under field conditions,
534 intraspecifically, zoeae of Pleocyemata species ingest a wide range of non-motile (and motile)
535 phytoplankton (and phytodetritus), from pico, through nano, to micro in size (Stickney &
536 Perkins 1981, Paul et al. 1989, Meyer-Harms & Harms 1993, Fileman et al. 2014). Such
537 indiscriminate ingestion of prey by those zoeae, including microzooplankton prey with limited
538 mobility used for the laboratory rearing, can be understood by widely known filter-feeding

539 habits of Pleocyemata larvae for any encountered prey of varying sizes (Rumrill et al. 1985,
540 Factor & Dexter 1993, Crain 1999, Kiørboe 2011, Wirtz 2012).

541 The indiscriminate filter-feeding of decapod crustacean zoeae could support their
542 growth and survival under non-motile prey-rich conditions in the dark (Harvey & Epifanio
543 1997, Hinz et al. 2001). In the water column of Amakusa-nada in the summer of 2006, most *N.*
544 *harmandi* zoeae stayed below 20 m depth, where photon flux density was less than 2% of that
545 at the surface, with the chl maximum layer at 22 to 24 m (Tamaki et al. 2010). In that water
546 column in August 2012, diatoms were the predominant taxonomic group, accounting for 57 to
547 81% of the total biovolume of the potential prey assemblage between 20 and 40 m water depths
548 (Fig. 6b). This tendency would be common in temperate coastal waters with higher nutrient
549 concentrations during the summertime (Harvey et al. 1997, Furuya et al. 2003). Therefore, if *N.*
550 *harmandi* zoeae indiscriminately ingest non-motile plankton by filter feeding in
551 Amakusa-nada, diatoms and their sinking detritus would be the primary diet.

552 Although SI analysis can be used to provide strong evidence for the trophic status of
553 meroplanktonic larvae of decapod crustaceans in the field, actual studies are scarce and limited
554 to bulk analyses (Schwamborn et al. 1999, 2002). Of the present 2 TP estimates for *N.*
555 *harmandi* zoeae from Amakusa-nada based on the $\delta^{15}\text{N}$ of amino acids (TP_{AAs}), one was 2.1
556 when using the most common measure suitable for metazoan food-chain ($\text{TP}_{\text{Glu/Phe}}$), while the
557 other was 2.7 when using a measure incorporating a possible protistan grazing pathway
558 ($\text{TP}_{\text{Ala/Phe}}$). Combination of these 2 estimates suggest that contribution of metazoans to the diet
559 of those zoeae is minimum ($\text{TP}_{\text{Glu/Phe}} \approx 2$) and that the primary diet is a mixture of
560 phytoplankton and heterotrophs mainly including protozoa ($\text{TP}_{\text{Ala/Phe}} = 2$ to 3). The potentially
561 mixed prey consisting of phytoplankton and protists was also reported for some
562 suspension-feeding smaller holozooplankton based on the 2 TP_{AAs} (e.g. $\text{TP}_{\text{Glu/Phe}} \approx 2$ and
563 $\text{TP}_{\text{Ala/Phe}} = 2.5$ to 3.0 for copepodite-stage individuals of *Calanus pacificus* and juveniles of

564 *Euphausia pacifica* in Californian coastal waters; Décima et al. 2017); the body size of *N.*
565 *harmandi* zoeae (3 to 7 mm) is larger than that of these zooplankton. Protists constituted a part
566 of the diet for reared brachyuran larvae (Lehto et al. 1998, Hinz et al. 2001), and nano- and
567 pico-eukaryotes and ciliates were contained in the digestive tract contents of wild-caught
568 brachyuran and gebiidean larvae (Fileman et al. 2014). The setal structure and motion of
569 mouthpart appendages of *N. harmandi* zoeae are inefficient in collecting sparse tiny cells and
570 in grasping motile plankton such as free-living ciliates and dinoflagellates (Somiya et al. 2014,
571 R. Somiya unpubl. data). Generally, phytodetritus is regarded as a complex aggregate with
572 heterotrophs (Turner 2015), which can be large enough for decapod crustacean larvae to collect
573 as well as large diatoms. Thus, phytoplankton, diatoms in particular (Fig. 6), and their detritus
574 with associated heterotrophs including protozoa are estimated to be the most likely diet of *N.*
575 *harmandi* zoeae.

576 The bulk SI analysis provided another clue to support the estimated contribution of
577 phytodetritus to the *N. harmandi* larval diet. The bulk SI ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$) values for the
578 latest-stage zoeae (V and VI) in Amakusa-nada are expected to have been equilibrated with
579 those for their food sources, following the observation of the SIs in whole-body tissue of zoeae
580 fed cultured diatoms and their stabilization around Day 20 (Fig. 4). In fact, in the dual bulk SI
581 diagram comprising POM, zooplankton (only in 2012), and the zoeae, the SI values for the
582 zoeae I and II shifted to the lighter ones as they grew to the zoeae V and VI both in 2012 and
583 2013 (Fig. 7). The mean (\pm SD) $\delta^{15}\text{N}$ values for zoeae V and VI in 2012 ($5.7 \pm 0.2\text{‰}$, excluding
584 outliers) were similar to those of the smaller zooplankton ($5.5 \pm 0.2\text{‰}$), and even lighter than
585 those of the larger zooplankton ($6.3 \pm 0.3\text{‰}$) (Fig. 7a). This suggests that copepod-dominated
586 zooplankton (0.3 to 3.5 mm in size) were not the potential food source for the *N. harmandi*
587 zoeae. A similar configuration of microzooplankton (presumed primary consumers) and
588 decapod crustacean zoeae in the dual bulk SI diagram was found for a plankton assemblage in

589 the coastal ocean off a mangrove estuary in northeastern Brazil (Schwamborn et al. 1999,
590 2002). Unexpectedly in the present study, the mean $\delta^{15}\text{N}$ value for the POM ($5.7 \pm 0.5\text{‰}$) was
591 not far below but rather close to the $\delta^{15}\text{N}$ value for the zoeae V and VI. This similarity poses a
592 question on the status of POM (presumed phytoplankton) as the main food source for the zoeae,
593 considering isotopic discrimination across trophic steps. Firstly, POM is not composed of
594 phytoplankton only, but of a mixture of phytoplankton, phytodetritus, zooplankton debris, and
595 fecal pellets (Lee 2002, Volkman & Tanoue 2002). With POM being such a complex mixture,
596 the $\delta^{15}\text{N}$ value might be lighter for live phytoplankton per se than for POM. Secondly, even if
597 the main constituent of POM from the chl maximum layers is live phytoplankton, its degraded
598 form (i.e. phytodetritus including heterotrophs), with altered SI values, could be the principal
599 diet of the *N. harmandi* zoeae. These larvae might selectively feed on sinking aggregates
600 composed of degraded lumps of diatoms with heterotrophs, which could be more easily
601 collected and ingested than loose cells.

602 The analyses of the dual bulk SI diagrams indicated that phytodetritus was likely an
603 important food source for *N. harmandi* zoeae, based on the SI shifts in POM during
604 degradation and larval TEFs (Fig. 7). Those zoeae stay mainly at 30 to 50 m depths in
605 Amakusa-nada (Tamaki et al. 2010), where sinking phytodetritus originated from the upper chl
606 maximum layer (usually 10 to 20 m depths except for August, with 30 to 40 m depths) may be
607 abundant. A potential daily shift in the SI values of POM derived from that layer, with
608 microbial degradation, is estimated at -0.07 to -0.09‰ d^{-1} for $\delta^{13}\text{C}$ and -0.11 to -0.18‰ d^{-1}
609 for $\delta^{15}\text{N}$, based on the 2 incubation experiments (Fig. 5). These alteration rates are comparable
610 to those reported in Lehmann et al. (2002), in which diatom-dominated POM was incubated
611 under different redox conditions (-0.06 to -0.13‰ d^{-1} for $\delta^{13}\text{C}$ and -0.10 to -0.12‰ d^{-1} for
612 $\delta^{15}\text{N}$; calculated here from those for the initial 10 d of the original 111 d incubation). Assuming
613 that live phytoplankton at the chl maximum layer in Amakusa-nada sink for a distance of 25 m

614 in 7 d at an average speed of 3.5 m d⁻¹ (cf. Kriest & Oschlies 2008), with the corresponding
615 SI-value alterations ($\Delta\delta^{13}\text{C} = -0.5$ to -0.6‰ wk⁻¹; $\Delta\delta^{15}\text{N} = -0.8$ to -1.3‰ wk⁻¹), the range of
616 the SI values for the expected degraded POM will be located at the lighter position than that for
617 the measured POM with large variations. In 2012, the expected $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of the
618 food source for *N. harmandi* zoeae V and VI inclusive based on their whole-body (including
619 exoskeleton)-specific TEFs (2.0‰ for $\delta^{13}\text{C}$; 1.9‰ for $\delta^{15}\text{N}$; Fig. 4) lay close to the edge lines
620 of the larger SI-box for the expected degraded-POM (expected FS [food source]-1 in Fig. 7a).
621 In a previous companion rearing experiment for *N. harmandi* juveniles, Yokoyama et al.
622 (2005) reported that TEFs for the whole body (-1.7‰ for $\delta^{13}\text{C}$; 2.3‰ for $\delta^{15}\text{N}$) were lower
623 than those for the muscle, probably due to the effect of chitin in the exoskeleton; no lipid
624 removal was made as in the present study. Based on these specific TEFs, there was no
625 candidate organic matter for the zoeal food sources (expected FS-2). Thus, it appears
626 indispensable to obtain the TEFs that are specific to the zoeal whole body, with its exoskeleton
627 probably not so fully developed as in juvenile exoskeleton.

628 The configuration of POM, presumed degraded-POM, and *N. harmandi* zoeae in the
629 dual bulk SI diagram for 2013 was largely similar to that for 2012, though the $\delta^{13}\text{C}$ values for
630 zoeae V and VI inclusive in 2012 were lighter by about 1.5‰ (Fig. 7b). One reason for the
631 latter might be a higher contribution of terrestrial dissolved inorganic carbon to primary
632 production, with relatively lighter $\delta^{13}\text{C}$ values than those in seawater (e.g. Boutton 1991). The
633 cumulative rainfall recorded at a meteorological observatory in Kumamoto Prefecture (32.813°
634 N, 130.707° E), located near Ariake Sound, during the rainy season of Japan (June and July)
635 was 3 times higher in 2012 than in 2013 (1200 vs. 430 mm; Japan Meteorological Agency Past
636 Weather Data; www.data.jma.go.jp/obd/stats/etrn/index.php). The average discharge of the
637 nearby largest river (Chikugo River emptying into the innermost Ariake Sound, with a 35%
638 watershed area in the sound; Iyama 2007) in this season were also 3 times higher in 2012 (493

664 Pleocyemata than previously thought, especially in coastal oceans connected to estuaries with
665 high primary production. In this demonstration, a combined approach (rearing with
666 pure-cultured phytoplankton, field surveys of nano- to micro-sized plankton assemblage, TP
667 analysis based on AA $\delta^{15}\text{N}$, determination of species-specific TEFs, and bulk SI analysis)
668 proved useful to estimate diatom-dominated phytoplankton and their detritus with heterotrophs
669 including protozoa as the principal diet of meroplanktonic larvae. The possible trophic status of
670 those planktotrophic larvae as the consumer relying mainly on phytoplankton would be
671 relevant to such aspects as (1) bottom-up effects of primary producers on the survival of those
672 larvae (Thorson 1950, Stickney & Perkins 1981, Olson & Olson 1989, Shirley & Shirley 1989,
673 Kirby et al. 2008), (2) reciprocal impact from those larvae to primary producers (Fileman et al.
674 2014), (3) vertical migration of those larvae in relation to phytoplankton and phytodetritus
675 distribution in the water column and its consequent horizontal transport (Pearre 2003,
676 Woodson & McManus 2007, Tamaki et al. 2010), and (4) biogeography of plankto- and
677 lecitho-trophic larval-type distribution with chlorophyll concentration distribution in the global
678 sea surface (Thorson 1950, Marshall et al. 2012).

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943

944

945 Table 1. Partially valid phytoplanktonic food items for the development of reared larvae in
 946 groups of the suborder Pleocyemata

Food items	Groups in Pleocyemata	Reference(s)
Diatoms	Brachyurans and/or anomurans	Hartman & Letterman (1978), Harms & Seeger (1989), Paul et al. (1989), Harms et al. (1991, 1994), Konishi et al. (1997)
Diatoms	Carideans	Stickney & Perkins (1981), Emmerson (1985)
Diatoms and dinoflagellates	Brachyurans	Incze & Paul (1983), Meyer-Harms & Harms (1993)
Dinoflagellates	Brachyurans	Perez & Sulkin (2005), Burnett & Sulkin (2007), Shaber & Sulkin (2007)
Dinoflagellates	Brachyurans and/or anomurans	Hinz et al. (2001)
Dinoflagellates and green alga	Brachyurans	Sulkin et al. (1998)
Dinoflagellates, green alga, and seagrass detritus	Brachyurans	Lehto et al. (1998)
Haptophyte alga	Brachyurans and/or anomurans	Mascetti & Wehrtmann (1996)

947

948

949 Fig. 1. Study region and location of tidal flats along the shoreline in mid-western Kyushu,
950 Japan. Water depth isopleths every 10 m were made by contouring (Surfer[®]8; Golden
951 Software) point data provided by the Hydrographic and Oceanographic Department, Japan
952 Coast Guard. Tidal flat areas, including the Tomioka sandflat, are indicated in black. Stn A in
953 Amakusa-nada was regularly visited for water and plankton sampling

954

955 Fig. 2. Newly fertilized embryo-volume-frequency distribution and newly hatched zoea I-total
956 length-frequency distribution of *Nihonotrypaea harmandi* specimens collected from late
957 August to early September 2014. The 2 normal distributions were fitted to each frequency
958 distribution

959

960 Fig. 3. Daily occurrence of decapodids of *Nihonotrypaea harmandi* in the 2 rearing tanks
961 between 3 August (Day 26) and 17 August (Day 40) 2013. Decapodids emerged first on Day
962 26. The 2 normal distributions were fitted to each frequency distribution

963

964 Fig. 4. Shift in (a) $\delta^{13}\text{C}$ and (b) $\delta^{15}\text{N}$ values (solid and blank circle plots, respectively) for the
965 whole-body tissue of *Nihonotrypaea harmandi* zoeae and decapodids in relation to that for
966 food (diatom, *Chaetoceros gracilis*: crosses) during the feeding experiment. See text for
967 sample numbers (n_s). The samples retrieved on Day 40 for stable isotope analysis consisted of
968 zoeae V and VI only. Dotted line and broken line in each panel: mean values for *N. harmandi*
969 and *C. gracilis*, respectively. $\Delta\delta^{13}\text{C}$ and $\Delta\delta^{15}\text{N}$ designate the trophic enrichment factors (TEFs)

970

971 Fig. 5. Shift in range or mean (\pm SD) $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values for particulate organic matter (POM)
972 with microbial degradation under dark conditions in the laboratory in June and July 2013. See
973 text for sample numbers (n_s)

974

975 Fig. 6. Vertical profiles of (a) abundance and (b) biovolume of nano- and micro-sized plankton
976 and their taxonomic compositions at Stn A in Amakusa-nada off mid-western Kyushu on 7
977 August 2012. Dt: diatom; Df: dinoflagellate; C: ciliate; O: other group including filamentous
978 cyanobacteria and unidentified organisms

979

980 Fig. 7. Dual bulk stable isotope (SI) plots for particulate organic matter (POM), zooplankton,
981 and zoeae I–VI of *Nihonotrypaea harmandi* collected from Stn A in Amakusa-nada off
982 mid-western Kyushu in (a) 2012 and (b) 2013. Samples from multiple cruises during the
983 summer were compiled for each year. Mean values for 2 successive zoeal stages inclusive (I
984 and II, III and IV) or the single mean values for the zoeae V and VI for *N. harmandi* and the
985 mean (\pm SD) values for the other components are indicated (see text for sample numbers, n_s).
986 Star: mean values for zoeae V and VI inclusive (SDs are omitted for simplicity; see text for
987 values). SI values for zooplankton in 2012 were distinctly lighter for the smaller 2 fractions
988 (0.3–0.8 and 0.8–1.2 mm in size) than for the larger 4 fractions (>1.2 mm), so that the
989 zooplankton were separated into 2 groups at 1.2 mm. The mean SI values of the 2 expected
990 food sources (FSs) corresponding to the mean values of the *N. harmandi* zoeae V and VI
991 inclusive were calculated from respective trophic enrichment factors (TEFs): FS-1 from zoeal
992 whole body-specific TEFs ($\Delta\delta^{13}\text{C} = 2.0\text{‰}$; $\Delta\delta^{15}\text{N} = 1.9\text{‰}$; Fig. 4) and FS-2 from *N. harmandi*
993 juvenile whole body-specific TEFs, respectively (based on Fig. 5 in Yokoyama et al. 2005).
994 Dark-shaded smaller box (rectangular area) and light-shaded larger box: expected smaller and
995 larger ranges of SIs for the presumed degraded-POM, respectively, derived from the mean SIs
996 for the POM and its SDs, respectively. The positions of 2 opposite corner points on the smaller
997 box are indicated by 2 arrows, each subjected to microbial degradation of the POM at the
998 corresponding daily decreasing rate in its mean SI values for 1 wk (Fig. 6)

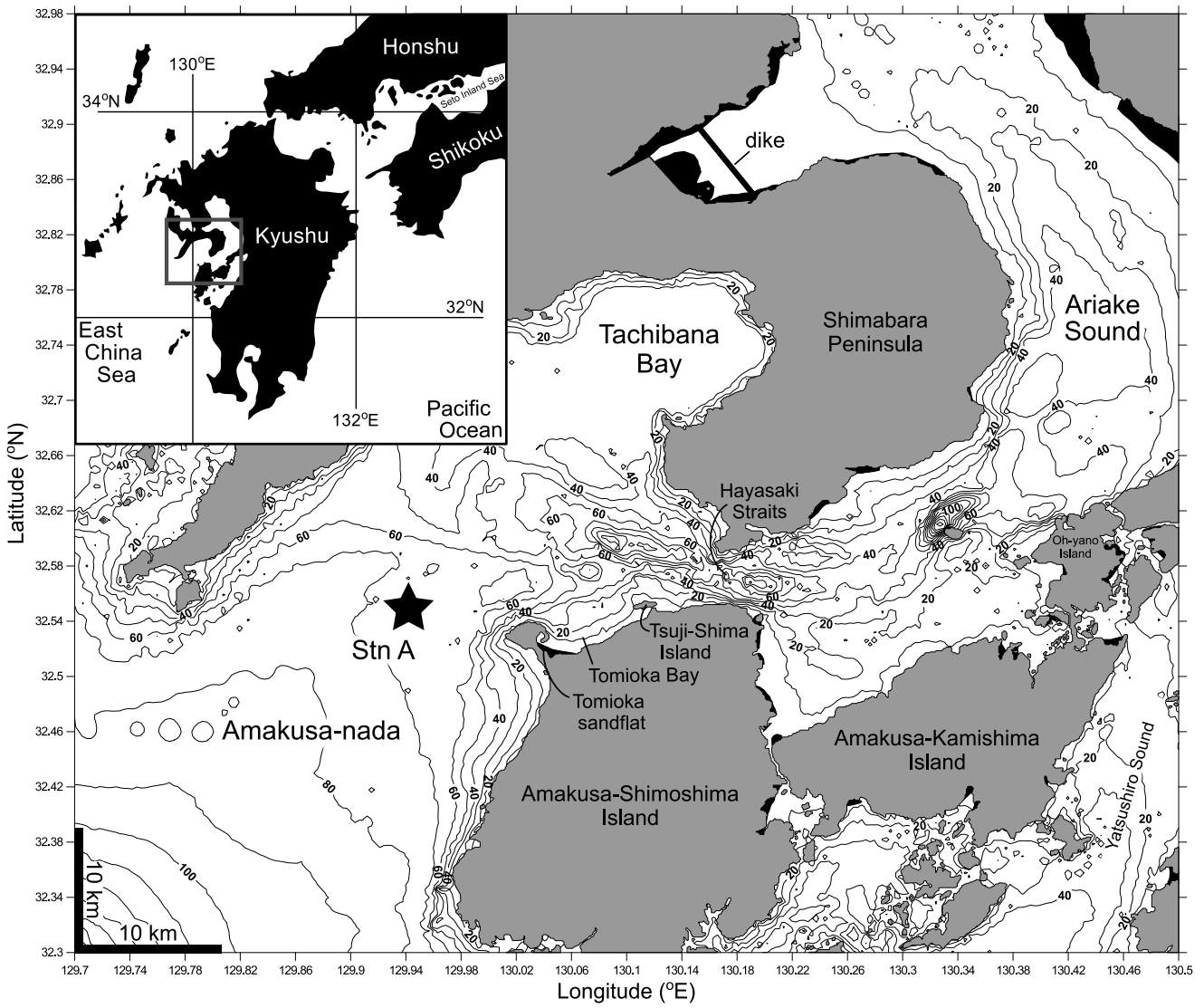


Fig. 1 (Umezawa et al., revised)

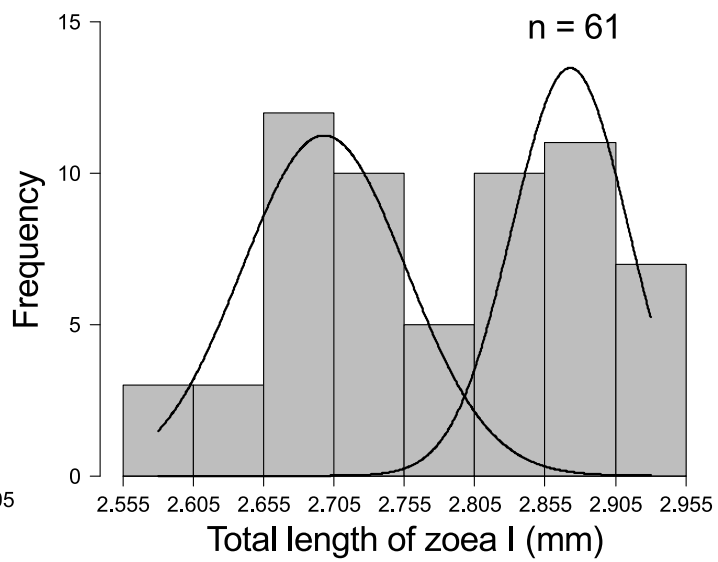
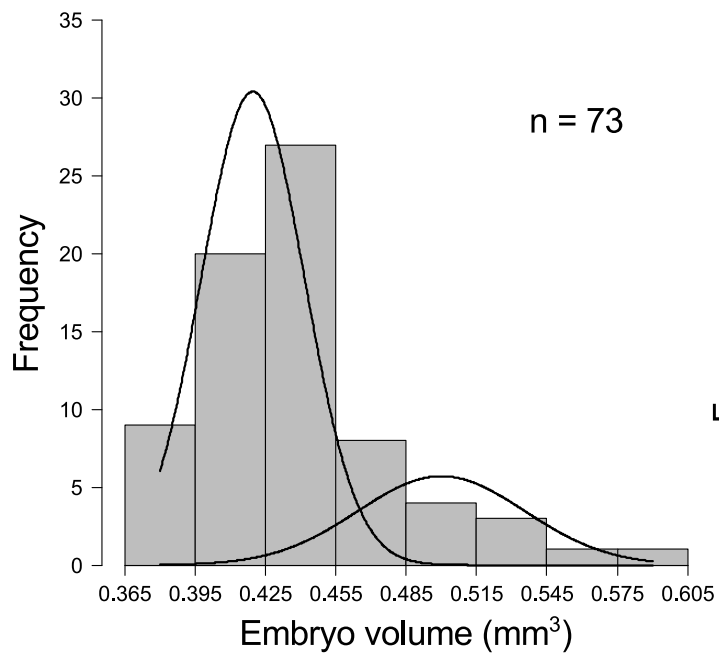


Fig. 2 (Umezawa et al., revised)

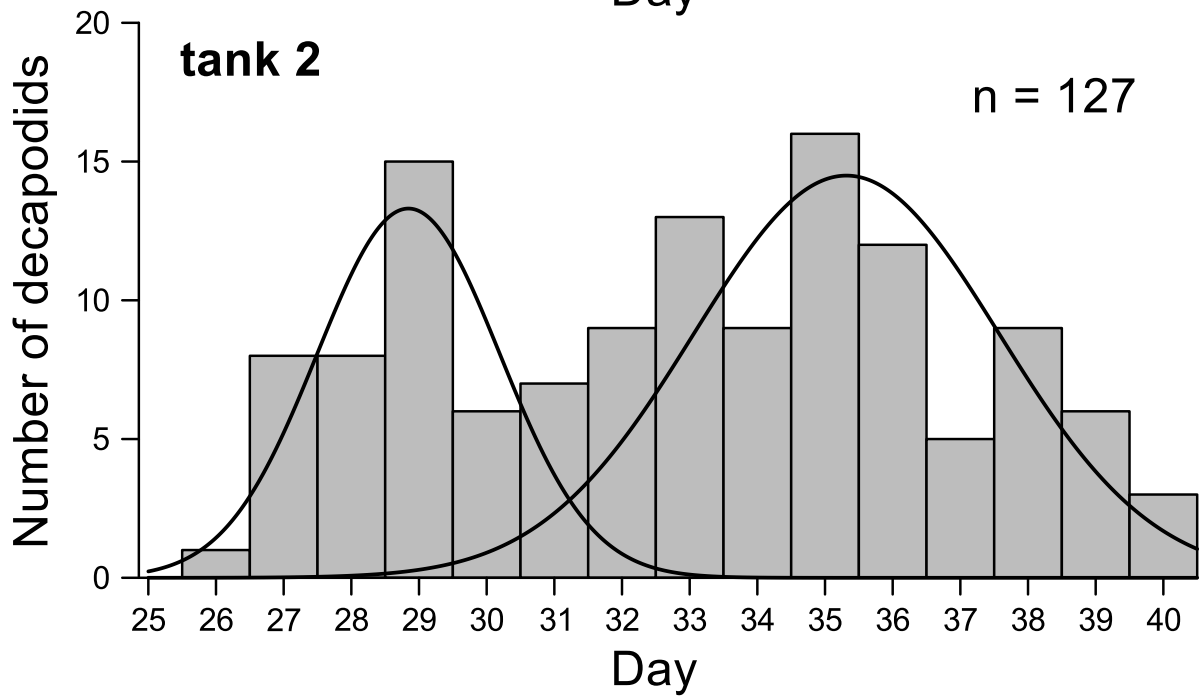
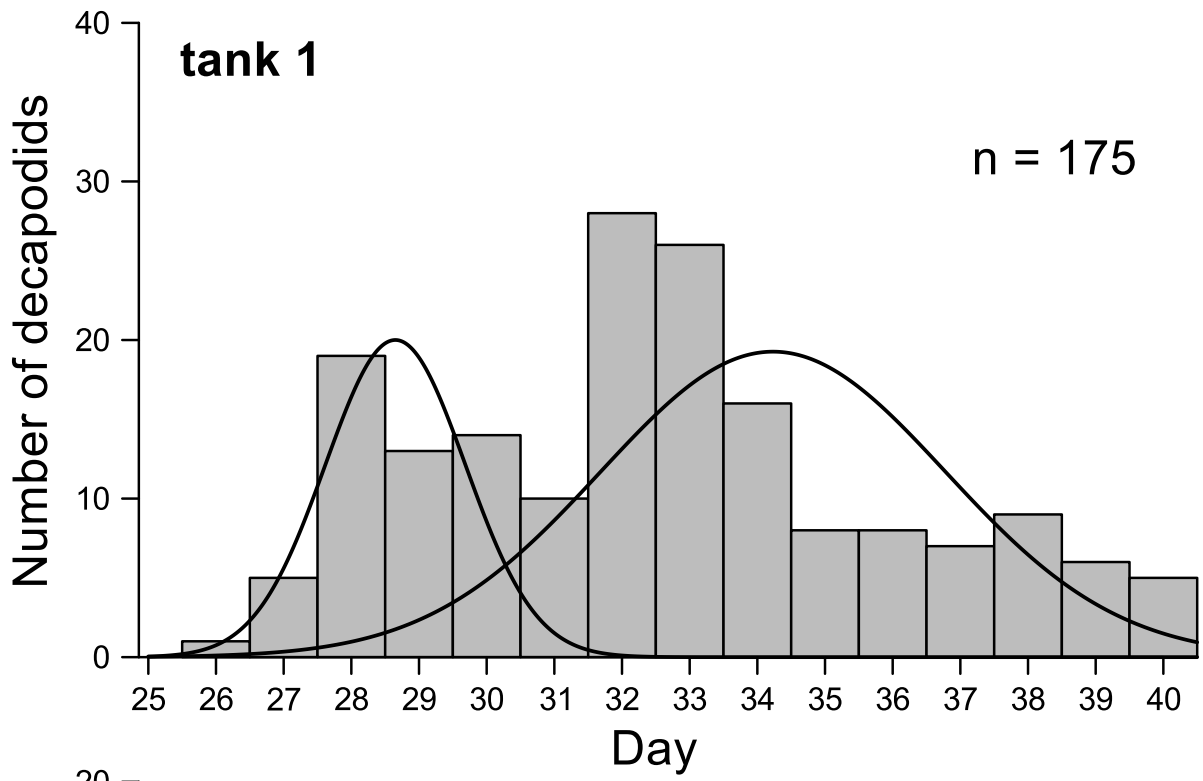


Fig. 3 (Umezawa et al., revised)

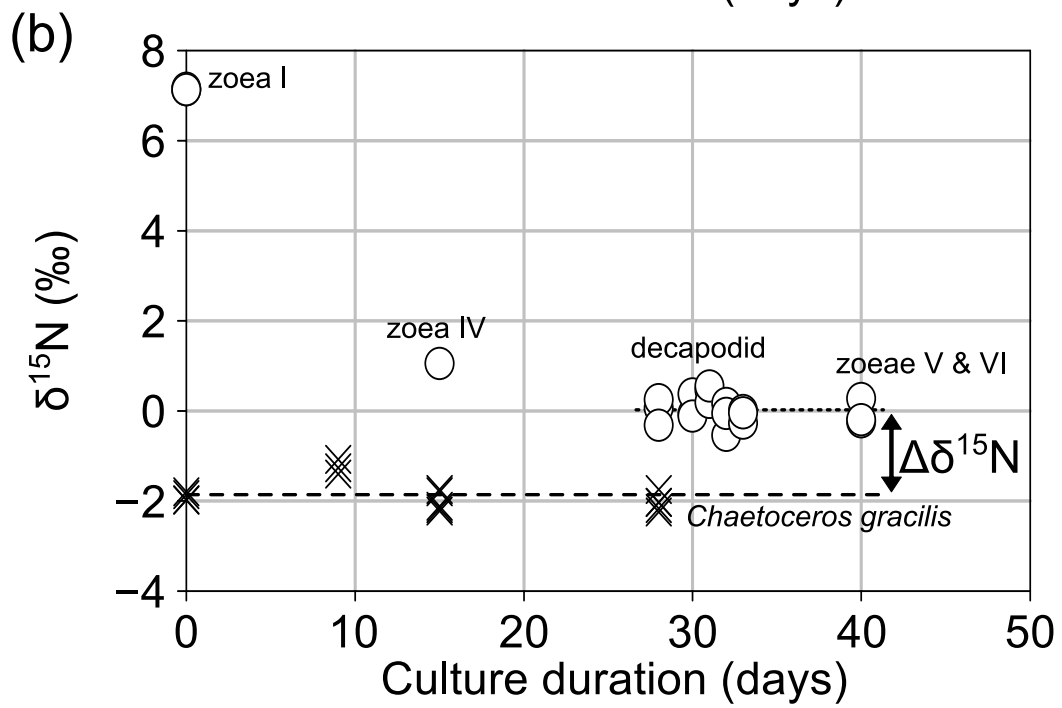
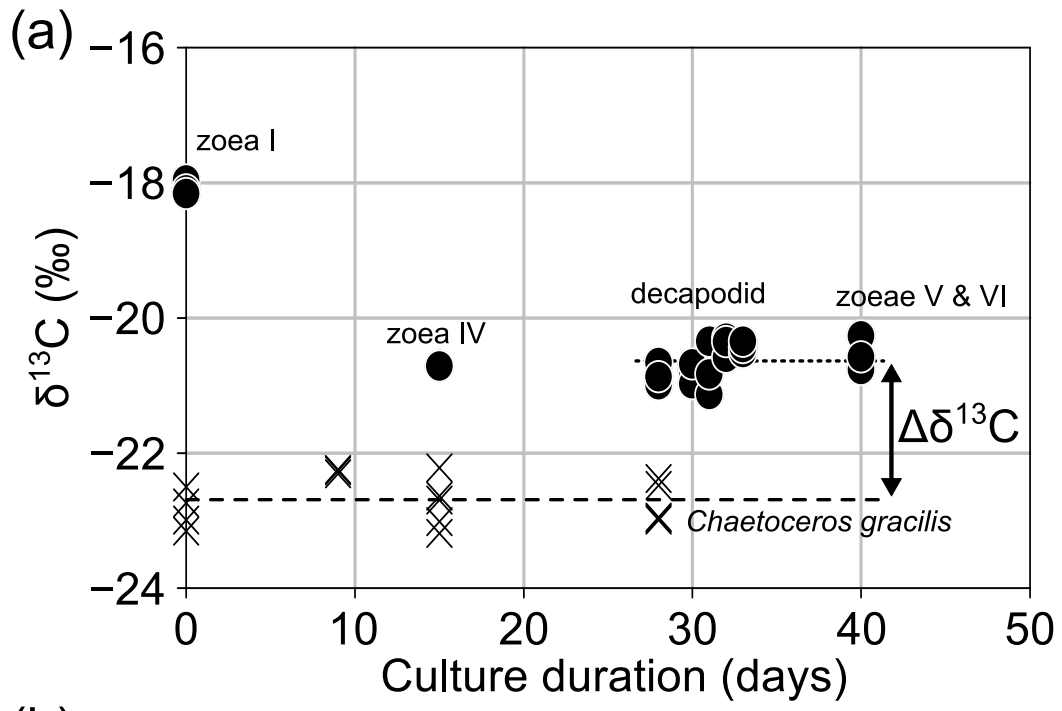


Fig. 4 (Umezawa et al., revised)

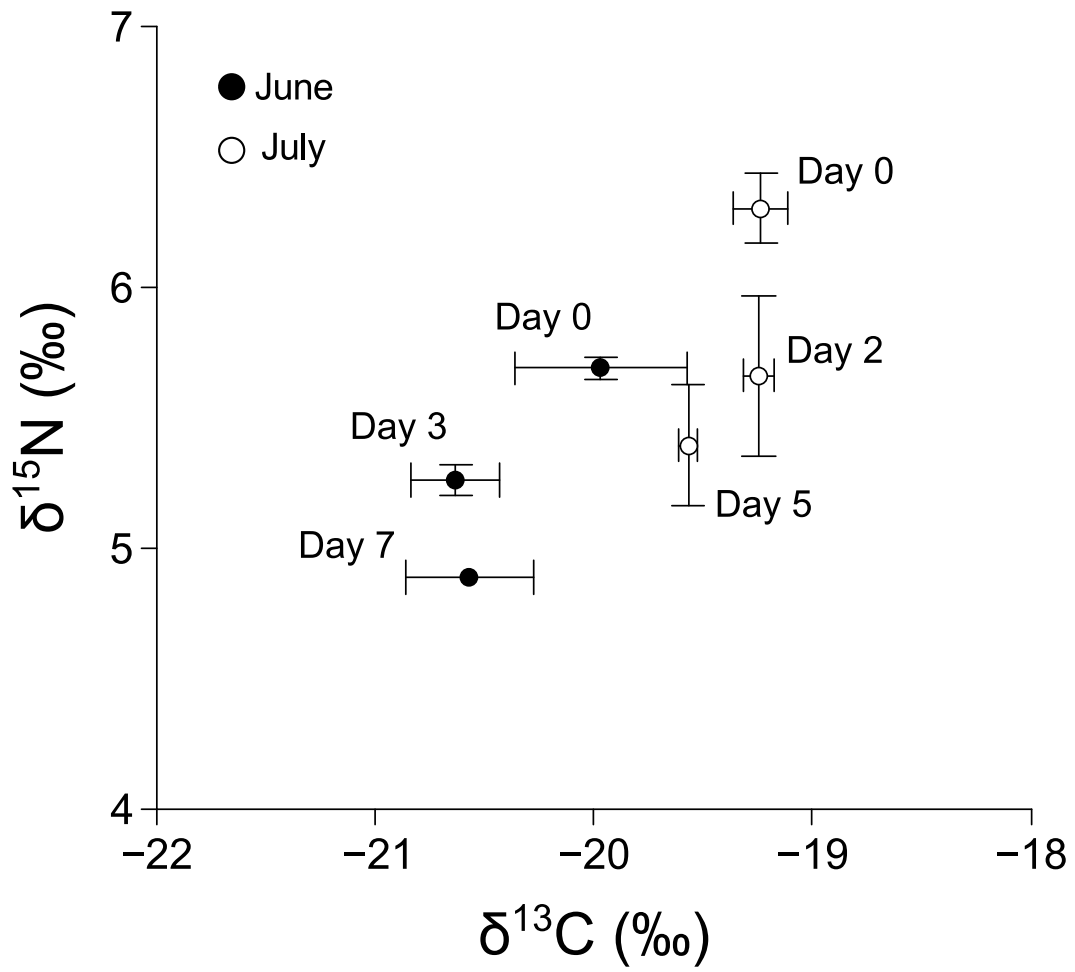


Fig. 5 (Umezawa et al., revised)

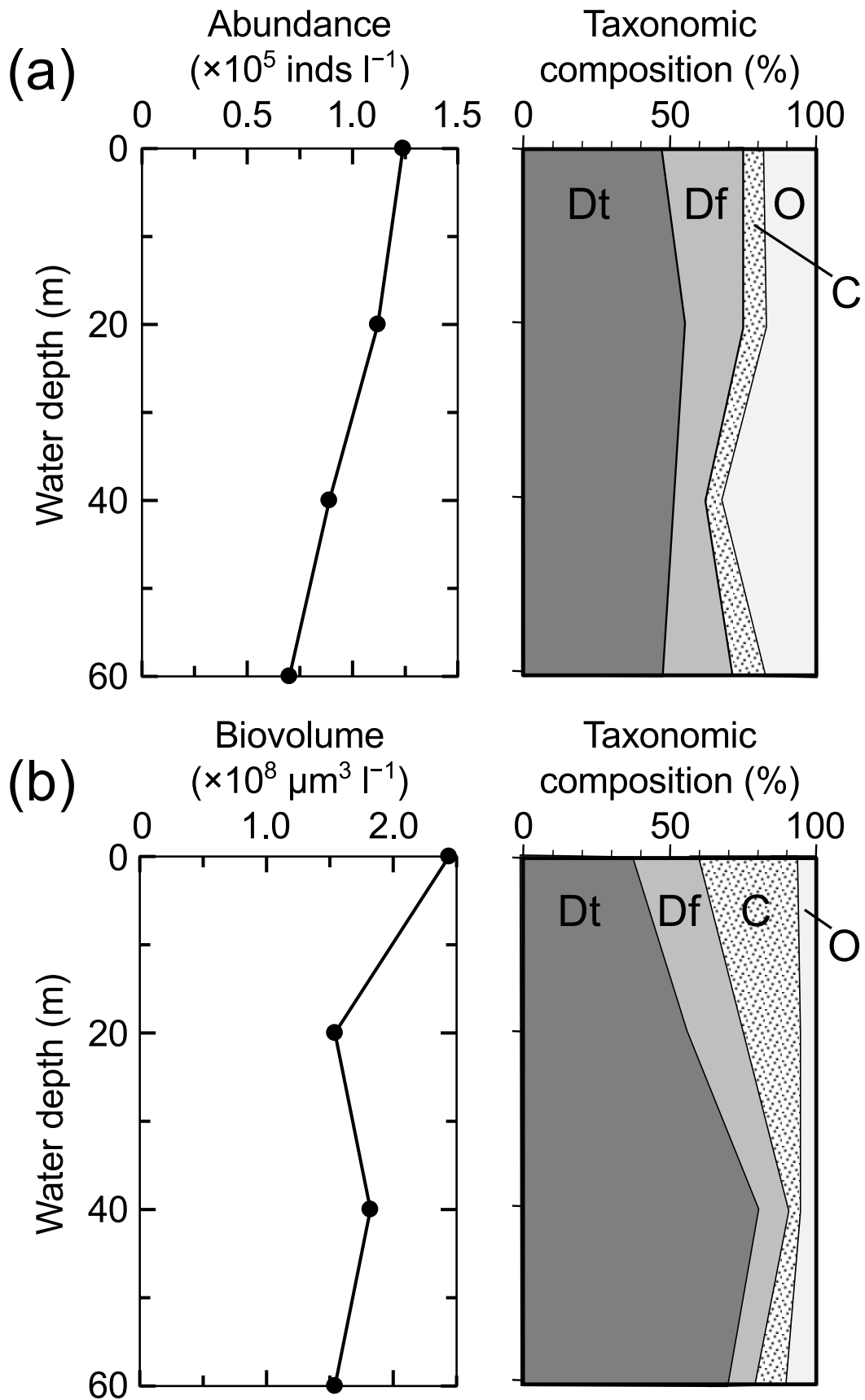


Fig. 6 (Umezawa et al., revised)

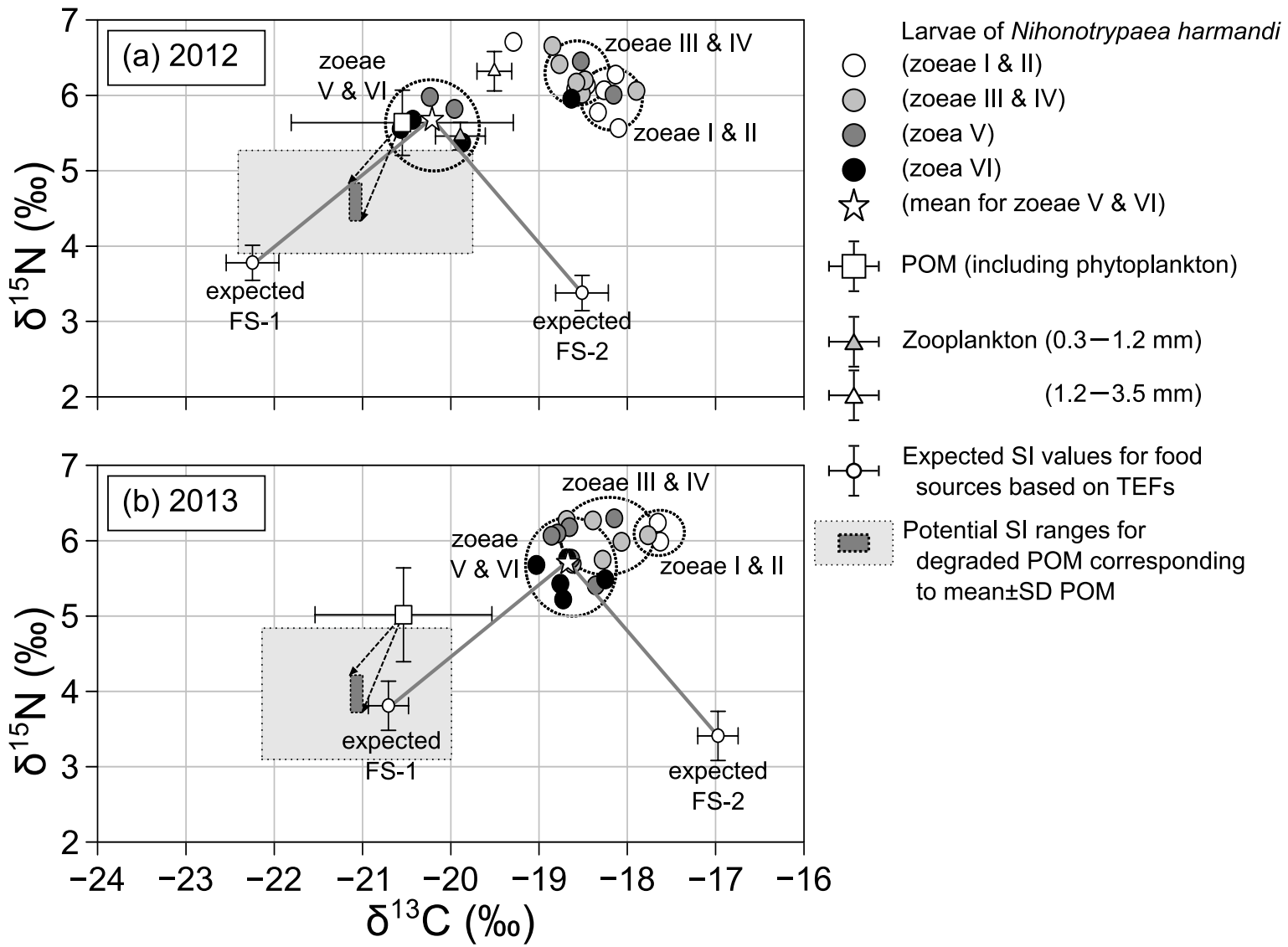


Fig. 7 (Umezawa et al., revised)